

FIGURE 1

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1 AAGTGCCTAAAGCTCTCCGAGGCCAAGGCCCTGCTACTGCCCGCGCTGCTTCTTAGTCCCGGTTCCGCGCTGGGTTCACCGCGCCGCGCGAGGA 100
 101 AGCCACTGCAACAGGACCCGAGTGGAGCGCGCGCAGCATGAAGCGCGCAGGCGCGCTCCATAGCGACGTCGGGACGGTCCGGCGCGCGCGGGGA 200
 201 AGGAAAAATGCAACATGCGCAGCAATGCAACAGAACAGCTGGGTGTTGAGATATTTGAACTGCGGACTGTGAGGAGAAATATTGAATCAGAGGATCGG 300
 M A A A M E T E Q L G V E I F E T A D C E E N I E S Q D R 29
 301 CCTAAATGGAGCCCTTTTATGTTGAGCGGTATCTCTGAGTCACTTAAAGCTGCTGTCGGATACCAAGAAATATCATGGCTACATGATGGCTAAGG 400
 30 P K L E P F F Y V E R Y S W S Q L K K L L A D T R K Y H G Y M K A 63
 401 CACCACATGATTTCATGTTTGTGAAGAGGAATGATCCAGATGCGACCTCATTCAGACAGAACTATTACCTTGCATGCTGCTGAGAACAGAGAAATAC 500
 63 P H D F H F V K R N D P D G P H S D R I Y Y L A M S G E N R E N T 96
 501 ACTGTTTTATTCGAAATTCGCAAACTATCAATAGAGCAGCAGCTCTTAACTGCTCTTGGAAAGCTCTTTTGGATCTTTTCAGGCAACACTGGACTAT 600
 96 L F Y S E I P K T I N R A A V L H L S W K P L L D L F Q A T L D Y 129
 601 CGAATGATATTCGAGAGGAAGAACTATTAAGAGAAAGAAACCGATTGGAACTGGAATGCTTCTTACGATTATCACCAAGGAAGTGGAACTTTTC 700
 130 G H Y S R E E L L R E R K R I G T V G I A S Y D Y H Q S G T F L 163
 701 TGTTCAGCGCGGTAGTGGAAATTTATCACTTAAAGATGGAGGCCCAAGGATTTACGCAACACCTTTAAGGCCCAATCTAGTGGAACTAGTTGCTC 800
 163 F Q A G S G I Y H V K D G G P Q G F T Q O P L R P N L V E T S C P 196
 801 CAACATACGGATGGATCCAAATTTATGCCCGCTGATCCAGACTGGATTGCTTTTATACATAGCAACGATATTTGGATATCAATCGTAACAGAGAA 900
 196 N I R M D P K L C P A D P D W I I H S N D I W I S N I V T R E 229
 901 GAAAGGAGACTCACTTATGTCACAAATGAGCTAGCCAACTGGAAGAGATGCGAGATCAGCTGGAGTGGCTACCTTTGTTCTCAAGAGAAATTTGATA 1000
 230 E R R L T Y V H N E L A N H E E D A R S A G V A T F V L Q E E F D R 263
 1001 GATATTTCTGGCTATTTGTTGTTGTCGAAAGCTGAAACAACTCCCACTGGTGGTAAAAATCTTAGAATTTCTATATGAAGAAATGATGAATCTGAGGTGGA 1100
 263 Y S G Y W W C P K A E T T P S G G K I L R I L Y E E N D E S E V E 296
 1101 AATTATTCATGTTACATCCCTTATGTTGGAAGCAAGGAGGCGAGATTCATCCGTTATCTTAAACAGGTACAGCAATCTTAAAGTCACTTTTAAAGATG 1200
 296 I I H V T S P K L E T R R A D S F R Y P K T G T A N P K V T F K H 329
 1201 TCAGAAATATGATTGATGCTGAGGAAGGATCATAGATGTCATAGATAAGGAATTAATCAACCTTTTGAAGATTCTATTTGAAGGAGTTGAATATATTG 1300
 330 S E I M I D A E G R I I D V I D K E L I Q P F E I L F E G V E Y I A 363
 1301 CCAGAGCTGATGAGCTCTGAGGGAATAATGCTTGGTCCATCTACTAGATGCTCCAGAGCTGGCTACAGATAGTCTGATCTCACTGAAATATT 1400
 363 K A G W T P E G K Y A W S I L L D R S O T R L Q I V L I S P E L 396
 1401 TATCCAGTAGAAGATGATGTTATGGAAGGCGAGAGCTATTGAGTCACTGCTGATTTCTGAGCGGCACTAATTATCTATGAAGAAACAGACATC 1500
 396 I P V E D D V M E R Q R L I E S V P D S V T P L I I Y E E T T D I 429
 1501 TGGATAAATATCCATGACATCTTTTATGTTTTCGCAAGTACAGAGAGGAAATGAGTTTATTTTTCCTCTGAAATGCAAAACAGGTTTTCCTCAT 1600
 430 W I N I H D I F H V F P Q S H E E E I E F I F A S E C K T G F R H L 463
 1601 TATACAAAATATCATCTATTTTAAAGGAAGCAATATAAAGCCAGTGGTGGGCTGCTGCTCAAGTGAATTCAGTGTCTATCAAGAGGAGAT 1700
 463 Y K I T S I L K E S K Y K R S S G G L P A P S D F K C P I K E E I 496
 1701 AGCAATTACCACTGGTGAATGCGAAGTCTTGGCGCGCATGGAATCAATATCAAGTTGATGAAGTCAGAGGCTGGTATATTTGAAGGCCACCAAGAC 1800
 496 A I T S G E W E V L G R H G S N I Q V D E V R R L V Y F E G T K D 529
 1801 TCCCTTTAGAGCATCACTGTACGTAGTCACTTACGTAATCTCGAGAGGTGACAGGCTGACTGACCGTGGCTACTCACAATCTTCTGCTCACTG 1900
 530 S P L E H H L Y V V S Y V N P G E V T R L T D R G Y S H C C I S Q 563
 1901 AGCACTGCACTCTTTTATAGTAAGTATAGTAACCAAGAAATCCACATCTGCTGCTGCTTACAGCTATCAAGTCTGAGATGACCACTTGCAT 2000
 563 H C D F F I S K Y S N Q K N P H C V S L Y K L S S P E D D P T C K 596
 2001 AACAAAGGAATTTTGGGCGACCATTTTGGATTGAGAGGCTGCTGCTGCTGACTATCTCTCCAGAAATTTTCTCTTTTGAAGTACTACTGATTTACA 2100
 596 T K E F W A T I L D S A G P L P D Y T P P E I F S F E S T T G F T 629
 2101 TTGATGGGATGCTCTACAAGCTCATGATCTACAGCTGGAAGAAATATCTACTGCTGCTTATATATGTTGCTCTCAAGTGCAGTTGTTGAATA 2200
 630 L Y G V L Y K P H D L Q P G K K Y F T V L F I Y G G P O V Q L V N N 663
 2201 ATGGTTTAAAGGAGTCAAGTATTTCCGCTTGAATACCTTAGCTCTTAGGTTATGCTGCTTGTAGTGAAGACACAGGGAATCTGTCACCGAGGCT 2300
 663 R P K G V K Y F R L N T L A S L G Y V V V V I D N R G S C H R G L 696
 2301 TAAATTTGAAGGCGCTTTTAAATATAAAATGGGTCAATAGAAATGAGCATCAGGTGGAAGGACTCCAAATATCTAGCTTCTGATATGATTTTCACTGAC 2400
 696 Y F E G A P F Y K M G Q I E I E D C V E G L Q Y L A S F Y D P I D 729
 2401 TTAGATGCTGTTGGCATCCAGGCTGCTCTATGGAAGGATACCTCTCCCTGATGGCATTAATGAGAGGTGAGATATCTTCAAGGTTGCTATTTCTGGG 2500
 710 I D P V G I H **25-57-25-100** G Y L S L M A L M O R S D I F R V A I A G A 763
 2501 CCCCAGTCACTCTGGAATCTTATGATACAGGATACACGGAAGCTTATATGGGTTATCTGACCAAGAAATGAAGGCTATTACTTAGGATCTGTCG 2600
 763 P V T L W I F Y D T G Y T E R Y H G H P D O N E Q G Y Y L G S V A 796
 2601 CATGCAAGCAGAAAGTTCCCTCTGAAACAAATGCTTTACTGCTCTTACATGCTTTCTGGATGAGAAATGCCATTTTGCACATACCAAGTATATTACTG 2700
 796 K Q A E K P P S E P K R L L L L H G F L **B** E N V H P A N T S I L 829
 2701 AGTTTTTATGAGGCGCTGGAAGCCATATGATTTACAGATCTATCTCCAGGAGAGACACAGCATAGAGTCTCTGAAATGCGGAGAACATTTGAAGTGC 2800
 830 S F L V R A G K P Y D L L Q I Y P Q E R **B** S I R V P E S G E H Y E L H 863
 2801 ATCTTTTGCACACTACCTTCAAGAAACCTTGGATCAGCTATTTCTGCTCTAAAGAGTATATAATTTTGACCTGTGTAGAACTCTGCTATACACTGGCTA 2900
 863 L L H Y L Q E N L G S R I A A L K V I * 882
 2901 TTTAACCAATGAGGAGGTTTAAATCAACAGAAACACAGAAATGATCATCATTTTGCATACCTGCCATGTAACATCTACTCTGAAAAATATAATGTTG 3000
 3001 CCAATGAGGGCTTACGGTTTGTGTTAGTAATCTAATACCTTAAACCCACATGCTCAAAAATCAATGATACATATCTCTGAGAGACCCAGCAATACCAT 3100
 3101 AGAATTACTAAAAAAAAAAAAAAAA 3127

FIGURE 2

[illegible]

FIGURE 3

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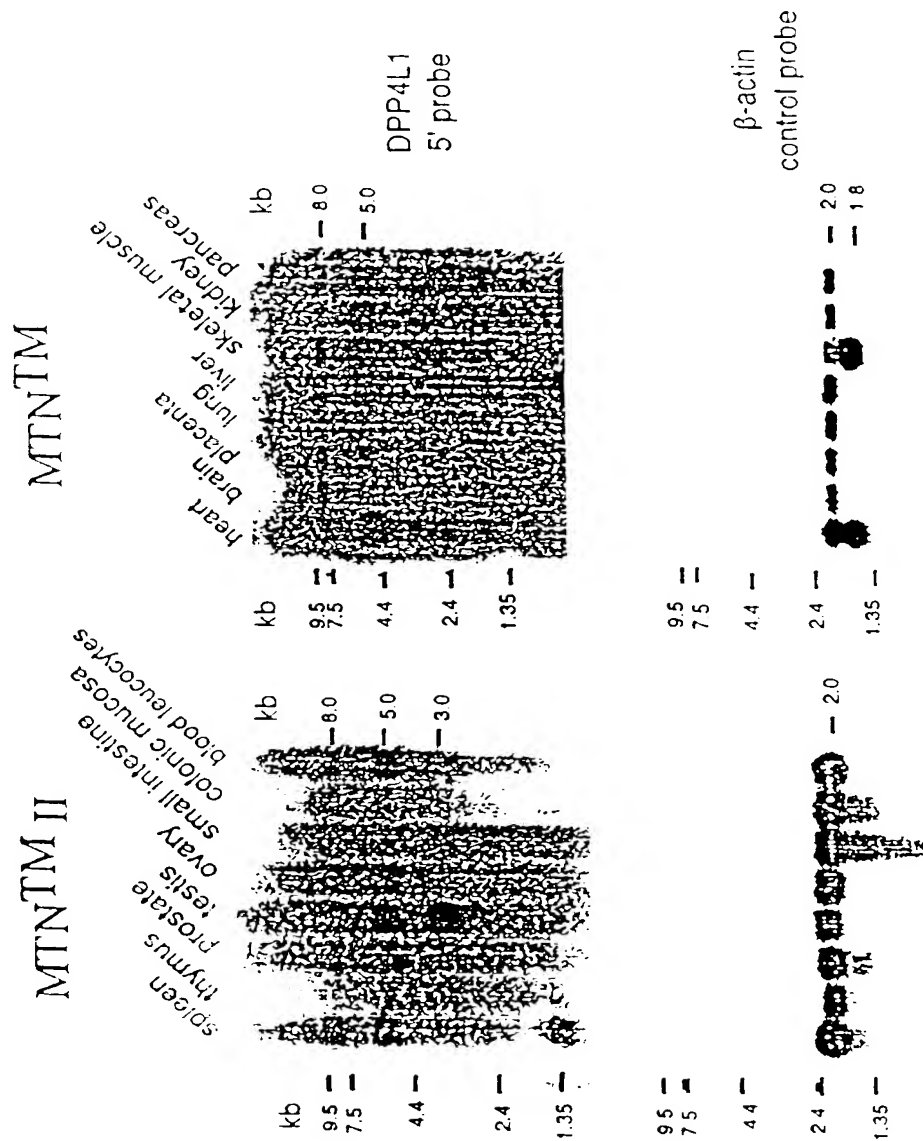


FIGURE 4

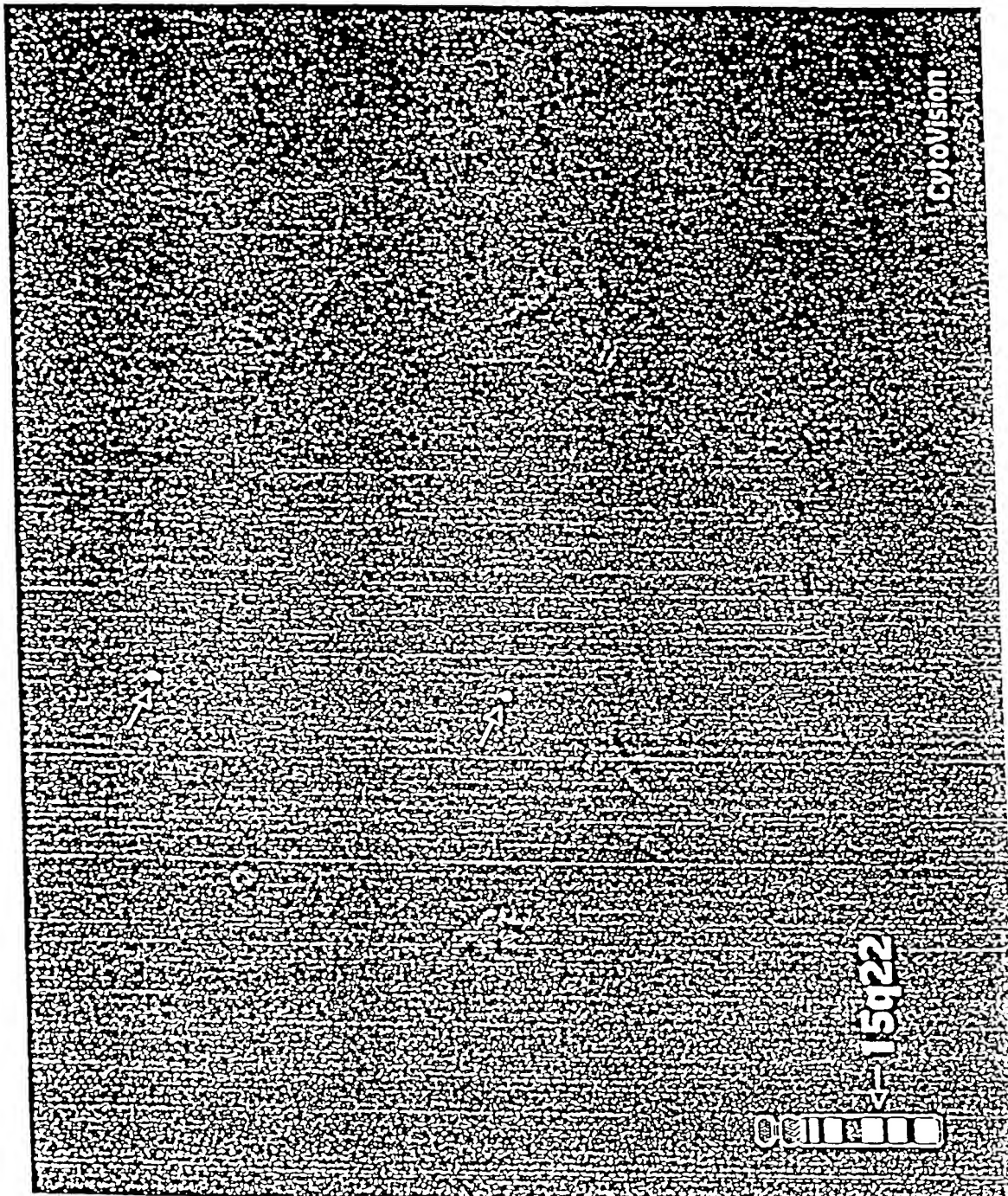
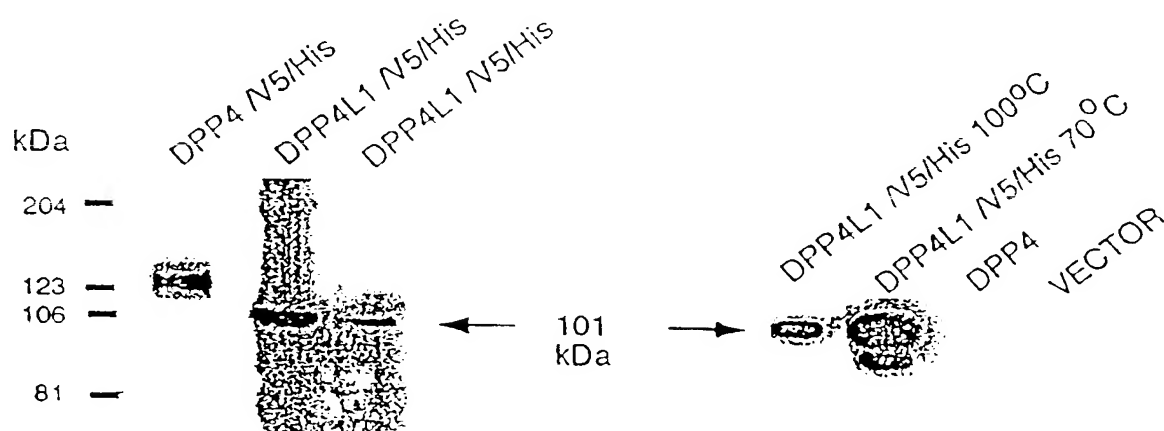


FIGURE 5

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FIGURE 6

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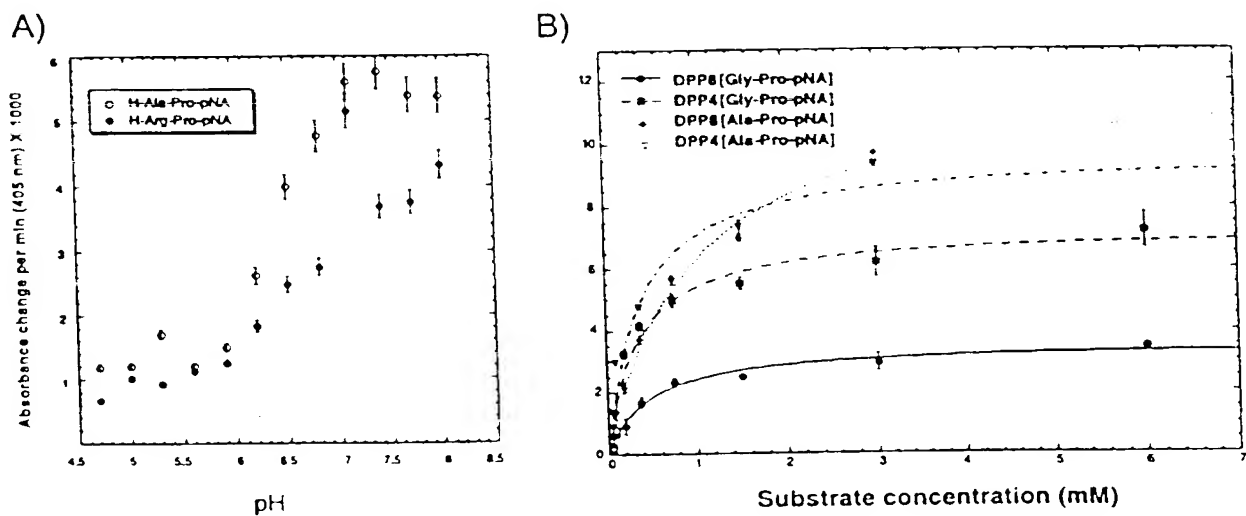


FIGURE 7

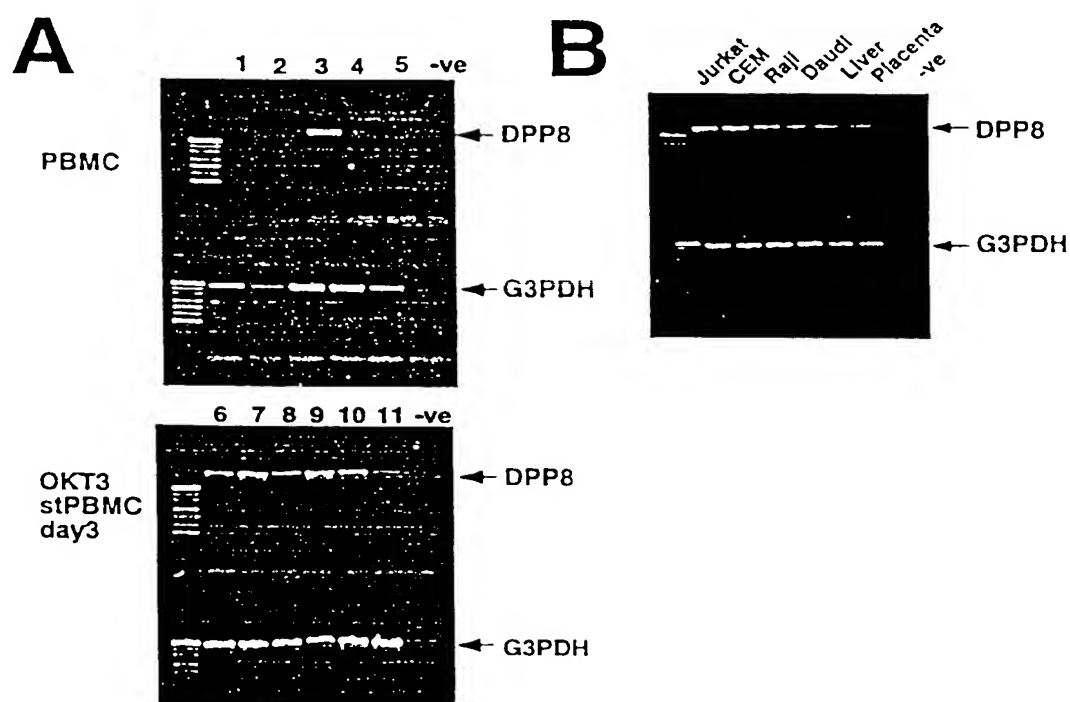


FIGURE 8

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FIGURE 9A

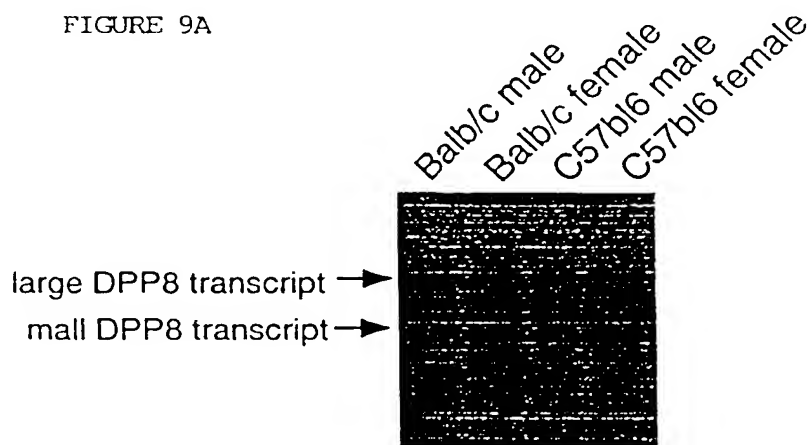


FIGURE 9B

